56818B~1 SEQUENCE LISTING

ĺ

<110> Washington University

(

$<\!\!120\!\!>\!\!$ Regulated Attenuation of Live Vaccines to Enhance Cross-Protective Immunogenicity

- <130> 56029-40434
- <140> PCT/US03/11802
- <141> 2003-04-15
- <150> US 60/373,626
- <151> 2002-04-18
- <150> US 60/372,616
- <151> 2002-04-15
- <160> 21
- <170> PatentIn version 3.1
- <210> 1
- <211> 25
- <212> DNA
- <213> Artificial Sequence
- <220>
- <223> Synthetic oligonucleotide
- <400> 1
- ggggtacccg aagacctgct gcgac

<210> 2

		56818B~1
<211>	25	300108~1
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	Synthetic oligonucleotide	
<400> 2 ccgaattcaa ttacacaca ccggt 25		
<210>	3	
<211>	25	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	Synthetic oligonucleotide	
<400> ccgaat	3 tcat cccgtcaggg aacgg	25
<210>		
<211>		
<212>		
<213>	Artificial Sequence	
<220>		
	Synthetic oligonucleotide	
<400>	-	
	tcat ttgccgctgc tggtc	25
<210>	5	
<211>	25	
<212>	DNA	
<213>	Artificial Sequence	

<220>

<223>	Synthetic oligonucleotide	
<400> ggggta	5 cctg gcaactttcc ggcgg	25
<210>	6	
<211>	25	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
	Synthetic oligonucleotide	
<400>	6 tcct tttatgacgc cggac	25
<210>	7	
<211>	25	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	Synthetic oligonucleotide	
<400> ccgaat	7 tctt atgattaagg aggca	25
<210>	8	
<211>	25	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	Synthetic oligonucleotide	
<400> gcgagc	8 tcgc gatatagttc gcata	25
∠210s	٥	

<211>	29	208188~1
<212>	DNA	
<213>	Artificial Sequence	
	·	
<220>		
<223>	Synthetic oligonucleotide	
<400>		29
yycyya	tcct gtcatctaat gagcggaat	25
<210>	10	
<211>	29	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	Synthetic oligonucleotide	
<400> gccgaa	10 ttca agtaacgata cctacaggc	29
<210>	11	
<211>	29	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
	Synthetic oligonucleotide	
<400> cgcgaa	11 ttca tcctacacgg caggtgaat	29
<210>	12	
<211>	29	
<212>	DNA	
<213>	Artificial Sequence	

<220>

<223>	Synthetic oligonucleotide	
<400> ccgaag	12 cttt cactgcaacc atgaatgac	29
<210>	13	
<211>	30	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	Synthetic oligonucleotide	
<400> cgggat	13 ccgt tatcggcaat ctggaggcaa	30
<210>	14	
<211>	30	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	Synthetic oligonucleotide	
<400> 14 catgcatgca ggcaggttca ggtacggtga 30		30
<210>	15	
<211>	27	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	Synthetic oligonucleotide	
<400> ggggta	15 ccta atcaacacta acagtct	27
<210s	16	

<211>	27	56818B~1	
<212>			
	Artificial Sequence		
1220	7.1		
<220>			
	Synthetic oligonucleotide		
<400>			
	tcag cagactgaac cgccagt		
<210>	17		
<211>	21		
<212>	DNA		
<213>	Artificial Sequence		
<220>			
<223>	Synthetic oligonucleotide		
<400>	17 ctcgg ggctttttca t		
ccyaac	ecegg ggeeceeca e		
<210>	18		
<211>	27		
<212>	DNA		
<213>	Artificial Sequence		
<220>			
<223>	Synthetic oligonucleotide		
<400>	18		
gcgagctctt caagaattgc cagagac			
<210>	19		
<211>	230		
<212>	PRT		

27

21

27

<400> 19

<213> Streptococcus pneumoniae

(

Leu Gln Ala Ser Asn Glu Ser Gln Arg Lys Glu Ala Asp Lys Lys Ile 1 10 15 Lys Glu Ala Thr Gln Arg Lys Asp Glu Ala Glu Ala Ala Phe Ala Thr 20 25 30 Ile Arg Thr Thr Ile Val Val Pro Glu Pro Ser Glu Leu Ala Glu Thr 35 40 45 Lys Lys Lys Ala Glu Glu Ala Thr Lys Glu Ala Glu Val Ala Lys Lys 50 60 Lys Ser Glu Glu Ala Ala Lys Glu Val Glu Val Glu Lys Asn Lys Ile 65 70 75 80 Leu Glu Gln Asp Ala Glu Asn Glu Lys Lys Ile Asp Val Leu Gln Asn 90 95 Lys Val Ala Asp Leu Glu Lys Gly Ile Ala Pro Tyr Gln Asn Glu Val 100 105 110 Ala Glu Leu Asn Lys Glu Ile Ala Arg Leu Gln Ser Asp Leu Lys Asp 115 120 125 Ala Glu Glu Asn Asn Val Glu Asp Tyr Ile Lys Glu Gly Leu Glu Gln 130 140 Ala Ile Thr Asn Lys Lys Ala Glu Leu Ala Thr Thr Gln Gln Asn Ile 145 150 155 160 Asp Lys Thr Gln Lys Asp Leu Glu Asp Ala Glu Leu Glu Leu Glu Lys 165 170 175 Val Leu Ala Thr Leu Asp Pro Glu Gly Lys Thr Gln Asp Glu Leu Asp 180 185 190 Lys Glu Ala Ala Glu Ala Glu Leu Asn Glu Lys Val Glu Ala Leu Gln 195 200 205 Asn Gln Val Ala Glu Leu Glu Glu Glu Leu Ser Lys Leu Glu Asp Asn 210 215 220 Leu Lys Asp Ala Glu Thr 225 230

<210> 20

<211> 257

<212> PRT

<213> Streptococcus pneumoniae

<400> 20

Leu Gln Ser Pro Val Ala Ser Gln Ser Lys Ala Glu Lys Asp Tyr Asp 1 10 15

Ala Ala Lys Lys Asp Ala Lys Asn Ala Lys Lys Ala Val Glu Asp Ala 20 25 30

Gln Lys Ala Leu Asp Asp Ala Lys Ala Ala Gln Lys Lys Tyr Asp Glu
35 40 45

Asp Gln Lys Lys Thr Glu Glu Lys Ala Ala Leu Glu Lys Ala Ala Ser 50 60

Glu Glu Met Asp Lys Ala Val Ala Ala Val Gln Gln Ala Tyr Leu Ala 65 70 75 80

Tyr Gln Gln Ala Thr Asp Lys Ala Ala Lys Asp Ala Ala Asp Lys Met
85 90 95

Ile Asp Glu Ala Lys Lys Arg Glu Glu Glu Ala Lys Thr Lys Phe Asn 100 105 110

Thr Val Arg Ala Met Val Val Pro Glu Pro Glu Gln Leu Ala Glu Thr 115 120 125

Lys Lys Lys Ser Glu Glu Ala Lys Gln Lys Ala Pro Glu Leu Thr Lys 130 140

Lys Leu Glu Glu Ala Lys Ala Lys Leu Glu Glu Ala Glu Lys Lys Ala 145 150 155 160

Thr Glu Ala Lys Gln Lys Val Asp Ala Glu Glu Val Ala Pro Gln Ala 165 170 175

Lys Ile Ala Glu Leu Glu Asn Gln Val His Arg Leu Glu Gln Glu Leu 180 185 190

Lys Glu Ile Asp Glu Ser Glu Ser Glu Asp Tyr Ala Lys Glu Gly Phe 195 200 205

Arg Ala Pro Leu Gln Ser Lys Leu Asp Ala Lys Lys Ala Lys Leu Ser Page 8

215

Lys Leu Glu Glu Leu Ser Asp Lys Ile Asp Glu Leu Asp Ala Glu Ile 225 230 235 240

Ala Lys Leu Glu Asp Gln Leu Lys Ala Ala Glu Glu Asn Asn Asn Val 245 250 255

Glu

<210> 21

210

<211> 487

<212> PRT

<213> Streptococcus pneumoniae

<400> 21

Leu Gln Ala Ser Asn Glu Ser Gln Arg Lys Glu Ala Asp Lys Lys Ile 1 10 15

Lys Glu Ala Thr Gln Arg Lys Asp Glu Ala Glu Ala Ala Phe Ala Thr 20 25 30

Ile Arg Thr Thr Ile Val Val Pro Glu Pro Ser Glu Leu Ala Glu Thr 35 40 45

Lys Lys Lys Ala Glu Glu Ala Thr Lys Glu Ala Glu Val Ala Lys Lys 50 60

Lys Ser Glu Glu Ala Ala Lys Glu Val Glu Val Glu Lys Asn Lys Ile 65 70 75 80

Leu Glu Gln Asp Ala Glu Asn Glu Lys Lys Ile Asp Val Leu Gln Asn 85 90 95

Lys Val Ala Asp Leu Glu Lys Gly Ile Ala Pro Tyr Gln Asn Glu Val 100 105 110

Ala Glu Leu Asn Lys Glu Ile Ala Arg Leu Gln Ser Asp Leu Lys Asp 115 120 125

Ala Glu Glu Asn Asn Val Glu Asp Tyr Ile Lys Glu Gly Leu Glu Gln 130 135 140 Ala Ile Thr Asn Lys Lys Ala Glu Leu Ala Thr Thr Gln Gln Asn Ile 145 150 155 160 Asp Lys Thr Gln Lys Asp Leu Glu Asp Ala Glu Leu Glu Leu Glu Lys
165 170 175 Val Leu Ala Thr Leu Asp Pro Glu Gly Lys Thr Gln Asp Glu Leu Asp 180 185 Lys Glu Ala Ala Glu Ala Glu Leu Asn Glu Lys Val Glu Ala Leu Gln 195 200 205 Asn Gln Val Ala Glu Leu Glu Glu Glu Leu Ser Lys Leu Glu Asp Asn 210 215 220 Leu Lys Asp Ala Glu Thr Leu Gln Ser Pro Val Ala Ser Gln Ser Lys 235 230 240 Ala Glu Lys Asp Tyr Asp Ala Ala Lys Lys Asp Ala Lys Asn Ala Lys 255 Lys Ala Val Glu Asp Ala Gln Lys Ala Leu Asp Asp Ala Lys Ala Ala 260 265 270 Gln Lys Lys Tyr Asp Glu Asp Gln Lys Lys Thr Glu Glu Lys Ala Ala 275 280 285 Leu Glu Lys Ala Ala Ser Glu Glu Met Asp Lys Ala Val Ala Ala Val 290 295 300 Gln Gln Ala Tyr Leu Ala Tyr Gln Gln Ala Thr Asp Lys Ala Ala Lys 305 310 315 Asp Ala Ala Asp Lys Met Ile Asp Glu Ala Lys Lys Arg Glu Glu Glu 325 330 335 Ala Lys Thr Lys Phe Asn Thr Val Arg Ala Met Val Val Pro Glu Pro 340 350 Glu Gln Leu Ala Glu Thr Lys Lys Lys Ser Glu Glu Ala Lys Gln Lys 355 360 365 Ala Pro Glu Leu Thr Lys Lys Leu Glu Glu Ala Lys Ala Lys Leu Glu 370 380 Glu Ala Glu Lys Lys Ala Thr Glu Ala Lys Gln Lys Val Asp Ala Glu 385 390 395 400

Glu Val Ala Pro Gln Ala Lys Ile Ala Glu Leu Glu Asn Gln Val His 405 410 415

Arg Leu Glu Gln Glu Leu Lys Glu Ile Asp Glu Ser Glu Ser Glu Asp 420 425 430

Tyr Ala Lys Glu Gly Phe Arg Ala Pro Leu Gln Ser Lys Leu Asp Ala 435 440 445

Lys Lys Ala Lys Leu Ser Lys Leu Glu Glu Leu Ser Asp Lys Ile Asp 450 460

Glu Leu Asp Ala Glu Ile Ala Lys Leu Glu Asp Gln Leu Lys Ala Ala 465 470 475 480

Glu Glu Asn Asn Asn Val Glu 485